



# SEQUENCE LISTING

<110> Amar, Salomon  
Tang, Xiaoren

<120> Novel LITAF Binding Site Peptides and Methods of Using the Same

<130> 50047/019002

<140> US 10/796,947

<141> 2004-03-10

<150> US 60/453,302

<151> 2003-03-10

<160> 33

<170> PatentIn version 3.3

<210> 1

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<212> PRT

<213> Homo sapiens

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Pro Ser Ala Pro Pro Ser Tyr Glu Glu Thr Val Ala Val Asn Ser Tyr  
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Tyr Pro Thr Pro Pro Ala Pro Met Pro Gly Pro Thr Thr Gly Leu Val  
35 40 45

Thr Gly Pro Asp Gly Lys Gly Met Asn Pro Pro Ser Tyr Tyr Thr Gln  
50 55 60

Pro Ala Pro Ile Pro Asn Asn Asn Pro Ile Thr Val Gln Thr Val Tyr  
65 70 75 80

Val Gln His Pro Ile Thr Phe Leu Asp Arg Pro Ile Gln Met Cys Cys  
85 90 95

Pro Ser Cys Asn Lys Met Ile Val Ser Gln Leu Ser Tyr Asn Ala Gly  
100 105 110

Ala Leu Thr Trp Leu Ser Cys Gly Ser Leu Cys Leu Leu Gly Val His  
115 120 125

Ser Gly Leu Leu Leu His Pro Leu Leu Arg Gly Cys Pro Ala Gly Arg  
 130 135 140

Gly Pro Leu Leu Ser Gln Leu Gln Ser Ser Pro Gly His Leu Gln Ala  
 145 150 155 160

Phe Val Gly Leu Ser Gln Thr Trp Arg Glu Pro Gly Ala Ala Gly Ser  
 165 170 175

Pro Phe His Leu Ser Ser Ser Phe Thr Pro Gly Gly Gly Ser Ala Leu  
 180 185 190

Val Val Ser Pro Leu Gln Gly Ala His Leu His Val Phe Phe Trp Gly  
 195 200 205

Glu Tyr Val Ala Lys Leu Thr Asn Leu Gln Thr Pro Glu Ile Ala Ala  
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Trp Ser Arg Ala  
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cctccccgcg gggtaaggcg ggcacccgc gagcgcaggg gtcctcttac tgctgatggc  
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accagctct gggcccagac gccgctcacc gtccaccgcc ggtgctgggt aaaatgtcgg  
 240

ttccaggacc ttaccaggcg gccactgggc cttcctcagc accatccgca cctccatcct  
 300

atgaagagac agtggctgtt aacagttatt accccacacc tccagctccc atgcttgggc  
 360

caactacggg gcttgtgacg gggcctgatg ggaagggcat gaatcctcct tcgtattata  
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cccagccagc gcccatcccc aataacaatc caattaccgt gcagacggtc tacgtgcagc  
480

accccatcac ctttttggac cgccctatcc aaatgtgttg tccttcctgc aacaagatga  
540

tcgtgagtca gctgtcctat aacgccgggtg ctctgacctg gctgtcctgc gggagcctgt  
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660

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ctgcttgag tcgtgcatag gacttgcaaa gacattcccc ttgagtgtca gttccacggc  
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tatcttcctg tgatctgcca tcagtggctc ttttttcctg cttccatggg cctttctggt  
1080

ggcagtctca aactgagaag ccacagttgc cttatTTTTTg aggtgtttct gccagagct  
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1200

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1260

cattagcaca cagattcaat ttaatttctt aatttttttt ttaaatacaa ggagggggct  
1320

attaacaccc agtacagaca tatccacaag gtcgtaaatg catgctagaa aaatagggct  
1380

ggatcttata actgccctgt ctccccttgt ttctctgtgc cagatcttca gtgccccttt  
1440

ccatacaggg atttttttct catagagtaa ttatatgaac agtttttatg acctcctttt  
1500

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1620

cccgaagtgc tgggattgca ggcataagct accatgctgg gcctgaacat aatttcaaga  
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Leu

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Val Gln Thr Val Tyr  
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<210> 32  
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Tyr Pro Tyr Asp Val Pro Asp Tyr Ala Ser Leu  
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cctca  
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